

# SEQUENCE LISTING

<110> Boehringer Ingelheim Vetmedica GmbH

<120> Recombinant attenuation of Porcine Reproductive and Respiratory Syndrome (PRRSV)

<130> 1-1120,25,26-1-C1

<140> to be assigned

<141> June 20, 2003

<150> 09/772,316

<151> January 26, 2001

<150> DE 10003371.7

<151> 2000-01-26

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<151> 2000-01-26

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<151> 2000-01-26

<160> 6

<170> PatentIn Ver. 2.1

<210> 1

<211> 2502

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 1

Met	Ser	Gly	Ile	Leu	Asp	Arg	Cys	Thr	Cys	Thr	Pro	Asn	Ala	Arg	Val
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Phe	Met	Ala	Glu	Gly	Gln	Val	Tyr	Cys	Thr	Arg	Cys	Leu	Ser	Ala	Arg
			20					25					30		
Ser	Leu	Leu	Pro	Leu	Asn	Leu	Gln	Val	Ser	Glu	Leu	Gly	Val	Leu	Gly
		35					40					45			
Leu	Phe	Tyr	Arg	Pro	Glu	Glu	Pro	Leu	Arg	Trp	Thr	Leu	Pro	Arg	Ala
	50					55					60				
Phe	Pro	Thr	Val	Glu	Cys	Ser	Pro	Ala	Gly	Ala	Cys	Trp	Leu	Ser	Ala
65					70					75					80
Ile	Phe	Pro	Ile	Ala	Arg	Met	Thr	Ser	Gly	Asn	Leu	Asn	Phe	Gln	Gln
				85					90					95	
Arg	Met	Val	Arg	Val	Ala	Ala	Glu	Leu	Tyr	Arg	Ala	Gly	Gln	Leu	Thr
			100					105					110		
Pro	Ala	Val	Leu	Lys	Ala	Leu	Gln	Val	Tyr	Glu	Arg	Gly	Cys	Arg	Trp
		115					120						125		

Tyr	Pro	Ile	Val	Gly	Pro	Val	Pro	Gly	Val	Ala	Val	Phe	Ala	Asn	Ser	130	135	140	
Leu	His	Val	Ser	Asp	Lys	Pro	Phe	Pro	Gly	Ala	Thr	His	Val	Leu	Thr	145	150	155	160
Asn	Leu	Pro	Leu	Pro	Gln	Arg	Pro	Lys	Pro	Glu	Asp	Phe	Cys	Pro	Phe	165	170	175	
Glu	Cys	Ala	Met	Ala	Thr	Val	Tyr	Asp	Ile	Gly	His	Asp	Ala	Val	Met	180	185	190	
Tyr	Val	Ala	Glu	Arg	Lys	Val	Ser	Trp	Ala	Pro	Arg	Gly	Gly	Asp	Glu	195	200	205	
Val	Lys	Phe	Glu	Ala	Val	Pro	Gly	Glu	Leu	Lys	Leu	Ile	Ala	Asn	Arg	210	215	220	
Leu	Arg	Thr	Ser	Phe	Pro	Pro	His	His	Thr	Val	Asp	Met	Ser	Lys	Phe	225	230	235	240
Ala	Phe	Thr	Ala	Pro	Gly	Cys	Gly	Val	Ser	Met	Arg	Val	Glu	Arg	Gln	245	250	255	
His	Gly	Cys	Leu	Pro	Ala	Asp	Thr	Val	Pro	Glu	Gly	Asn	Cys	Trp	Trp	260	265	270	
Ser	Leu	Phe	Asp	Leu	Leu	Pro	Leu	Glu	Val	Gln	Asn	Lys	Glu	Ile	Arg	275	280	285	
His	Ala	Asn	Gln	Phe	Gly	Tyr	Gln	Thr	Lys	His	Gly	Val	Ser	Gly	Lys	290	295	300	
Tyr	Leu	Gln	Arg	Arg	Leu	Gln	Val	Asn	Gly	Leu	Arg	Ala	Val	Thr	Asp	305	310	315	320
Leu	Asn	Gly	Pro	Ile	Val	Val	Gln	Tyr	Phe	Ser	Val	Lys	Glu	Ser	Trp	325	330	335	
Ile	Arg	His	Leu	Lys	Leu	Ala	Gly	Glu	Pro	Ser	Tyr	Ser	Gly	Phe	Glu	340	345	350	
Asp	Leu	Leu	Arg	Ile	Arg	Val	Glu	Pro	Asn	Thr	Ser	Pro	Leu	Ala	Asp	355	360	365	
Lys	Glu	Glu	Lys	Ile	Phe	Arg	Phe	Gly	Ser	His	Lys	Trp	Tyr	Gly	Ala	370	375	380	
Gly	Lys	Arg	Ala	Arg	Lys	Ala	Arg	Ser	Cys	Ala	Thr	Ala	Thr	Val	Ala	385	390	395	400
Gly	Arg	Ala	Leu	Ser	Val	Arg	Glu	Thr	Arg	Gln	Ala	Lys	Glu	His	Glu	405	410	415	
Val	Ala	Gly	Ala	Asn	Lys	Ala	Glu	His	Leu	Lys	His	Tyr	Ser	Pro	Pro	420	425	430	

Ala	Glu	Gly	Asn	Cys	Gly	Trp	His	Cys	Ile	Ser	Ala	Ile	Ala	Asn	Arg	435	440	445
Met	Val	Asn	Ser	Lys	Phe	Glu	Thr	Thr	Leu	Pro	Glu	Arg	Val	Arg	Pro	450	455	460
Pro	Asp	Asp	Trp	Ala	Thr	Asp	Glu	Asp	Leu	Val	Asn	Ala	Ile	Gln	Ile	465	470	475
Leu	Arg	Leu	Pro	Ala	Ala	Leu	Asp	Arg	Asn	Gly	Ala	Cys	Thr	Ser	Ala	485	490	495
Lys	Tyr	Val	Leu	Lys	Leu	Glu	Gly	Glu	His	Trp	Thr	Val	Thr	Val	Thr	500	505	510
Pro	Gly	Met	Ser	Pro	Ser	Leu	Leu	Pro	Leu	Glu	Cys	Val	Gln	Gly	Cys	515	520	525
Cys	Gly	His	Lys	Gly	Gly	Leu	Gly	Ser	Pro	Asp	Ala	Val	Glu	Val	Ser	530	535	540
Gly	Phe	Asp	Pro	Ala	Cys	Leu	Asp	Arg	Leu	Ala	Glu	Val	Met	His	Leu	545	550	555
Pro	Ser	Ser	Ala	Ile	Pro	Ala	Ala	Leu	Ala	Glu	Met	Ser	Gly	Asp	Ser	565	570	575
Asp	Arg	Ser	Ala	Ser	Pro	Val	Thr	Thr	Val	Trp	Thr	Val	Ser	Gln	Phe	580	585	590
Phe	Ala	Arg	His	Ser	Gly	Gly	Asn	His	Pro	Asp	Gln	Val	Arg	Leu	Gly	595	600	605
Lys	Ile	Ile	Ser	Leu	Cys	Gln	Val	Ile	Glu	Asp	Cys	Cys	Cys	Ser	Gln	610	615	620
Asn	Lys	Thr	Asn	Arg	Val	Thr	Pro	Glu	Glu	Val	Ala	Ala	Lys	Ile	Asp	625	630	635
Leu	Tyr	Leu	Arg	Gly	Ala	Thr	Asn	Leu	Glu	Glu	Cys	Leu	Ala	Arg	Leu	645	650	655
Glu	Lys	Ala	Arg	Pro	Pro	Arg	Val	Ile	Asp	Thr	Ser	Phe	Asp	Trp	Asp	660	665	670
Val	Val	Leu	Pro	Gly	Val	Glu	Ala	Ala	Thr	Gln	Thr	Ile	Lys	Leu	Pro	675	680	685
Gln	Val	Asn	Gln	Cys	Arg	Ala	Leu	Val	Pro	Val	Val	Thr	Gln	Lys	Ser	690	695	700
Leu	Asp	Asn	Asn	Ser	Val	Pro	Leu	Thr	Ala	Phe	Ser	Leu	Ala	Asn	Tyr	705	710	715
Tyr	Tyr	Arg	Ala	Gln	Gly	Asp	Glu	Val	Arg	His	Arg	Glu	Arg	Leu	Thr	725	730	735

Ala	Val	Leu	Ser	Lys	Leu	Glu	Lys	Val	Val	Arg	Glu	Glu	Tyr	Gly	Leu	
			740					745					750			
Met	Pro	Thr	Glu	Pro	Gly	Pro	Arg	Pro	Thr	Leu	Pro	Arg	Gly	Leu	Asp	
		755					760					765				
Glu	Leu	Lys	Asp	Gln	Met	Glu	Glu	Asp	Leu	Leu	Lys	Leu	Ala	Asn	Ala	
	770					775					780					
Gln	Thr	Thr	Ser	Asp	Met	Met	Ala	Trp	Ala	Val	Glu	Gln	Val	Asp	Leu	
785					790					795					800	
Lys	Thr	Trp	Val	Lys	Asn	Tyr	Pro	Arg	Trp	Thr	Pro	Pro	Pro	Pro	Pro	
				805					810					815		
Pro	Lys	Val	Gln	Pro	Arg	Lys	Thr	Lys	Pro	Val	Lys	Ser	Leu	Pro	Glu	
			820					825					830			
Arg	Lys	Pro	Val	Pro	Ala	Pro	Arg	Arg	Lys	Val	Gly	Ser	Asp	Cys	Gly	
		835					840					845				
Ser	Pro	Val	Ser	Leu	Gly	Gly	Asp	Val	Pro	Asn	Ser	Trp	Glu	Asp	Leu	
	850					855					860					
Ala	Val	Ser	Ser	Pro	Phe	Asp	Leu	Pro	Thr	Pro	Pro	Glu	Pro	Ala	Thr	
865					870					875					880	
Pro	Ser	Ser	Glu	Leu	Val	Ile	Val	Ser	Ser	Pro	Gln	Cys	Ile	Phe	Arg	
				885					890					895		
Pro	Ala	Thr	Pro	Leu	Ser	Glu	Pro	Ala	Pro	Ile	Pro	Ala	Pro	Arg	Gly	
			900					905					910			
Thr	Val	Ser	Arg	Pro	Val	Thr	Pro	Leu	Ser	Glu	Pro	Ile	Pro	Val	Pro	
		915					920						925			
Ala	Pro	Arg	Arg	Lys	Phe	Gln	Gln	Val	Lys	Arg	Leu	Ser	Ser	Ala	Ala	
	930					935					940					
Ala	Ile	Pro	Pro	Tyr	Gln	Asp	Glu	Pro	Leu	Asp	Leu	Ser	Ala	Ser	Ser	
945					950					955					960	
Gln	Thr	Glu	Tyr	Glu	Ala	Ser	Pro	Pro	Ala	Pro	Pro	Gln	Ser	Gly	Gly	
				965					970					975		
Val	Leu	Gly	Val	Glu	Gly	His	Glu	Ala	Glu	Glu	Thr	Leu	Ser	Glu	Ile	
			980					985					990			
Ser	Asp	Met	Ser	Gly	Asn	Ile	Lys	Pro	Ala	Ser	Val	Ser	Ser	Ser	Ser	
		995				1000						1005				
Ser	Leu	Ser	Ser	Val	Arg	Ile	Thr	Arg	Pro	Lys	Tyr	Ser	Ala	Gln	Ala	
	1010					1015					1020					
Ile	Ile	Asp	Ser	Gly	Gly	Pro	Cys	Ser	Gly	His	Leu	Gln	Glu	Val	Lys	
1025					1030					1035					1040	

Glu Thr Cys Leu Ser Val Met Arg Glu Ala Cys Asp Ala Thr Lys Leu		
1045	1050	1055
Asp Asp Pro Ala Thr Gln Glu Trp Leu Ser Arg Met Trp Asp Arg Val		
1060	1065	1070
Asp Met Leu Thr Trp Arg Asn Thr Ser Val Tyr Gln Ala Ile Cys Thr		
1075	1080	1085
Leu Asp Gly Arg Leu Lys Phe Leu Pro Lys Met Ile Leu Glu Thr Pro		
1090	1095	1100
Pro Pro Tyr Pro Cys Glu Phe Val Met Met Pro His Thr Pro Ala Pro		
1105	1110	1115
Ser Val Gly Ala Glu Ser Asp Leu Thr Ile Gly Ser Val Ala Thr Glu		
1125	1130	1135
Asp Val Pro Arg Ile Leu Glu Lys Ile Glu Asn Val Gly Glu Met Ala		
1140	1145	1150
Asn Gln Gly Pro Leu Ala Phe Ser Glu Asp Lys Pro Val Asp Asp Gln		
1155	1160	1165
Leu Val Asn Asp Pro Arg Ile Ser Ser Arg Arg Pro Asp Glu Ser Thr		
1170	1175	1180
Ser Ala Pro Ser Ala Gly Thr Gly Gly Ala Gly Ser Phe Thr Asp Leu		
1185	1190	1195
Pro Pro Ser Asp Gly Ala Asp Ala Asp Gly Gly Gly Pro Phe Arg Thr		
1205	1210	1215
Val Lys Arg Lys Ala Glu Arg Leu Phe Asp Gln Leu Ser Arg Gln Val		
1220	1225	1230
Phe Asp Leu Val Ser His Leu Pro Val Phe Phe Ser Arg Leu Phe Tyr		
1235	1240	1245
Pro Gly Gly Gly Tyr Ser Pro Gly Asp Trp Gly Phe Ala Ala Phe Thr		
1250	1255	1260
Leu Leu Cys Leu Phe Leu Cys Tyr Ser Tyr Pro Ala Phe Gly Ile Ala		
1265	1270	1275
Pro Leu Leu Gly Val Phe Ser Gly Ser Ser Arg Arg Val Arg Met Gly		
1285	1290	1295
Val Phe Gly Cys Trp Leu Ala Phe Ala Val Gly Leu Phe Lys Pro Val		
1300	1305	1310
Ser Asp Pro Val Gly Ala Ala Cys Glu Phe Asp Ser Pro Glu Cys Arg		
1315	1320	1325
Asn Ile Leu His Ser Phe Glu Leu Leu Lys Pro Trp Asp Pro Val Arg		
1330	1335	1340

Ser Leu Val Val Gly Pro Val Gly Leu Gly Leu Ala Ile Leu Gly Arg  
 1345 1350 1355 1360  
 Leu Leu Gly Gly Ala Arg Cys Ile Trp His Phe Leu Leu Arg Leu Gly  
 1365 1370 1375  
 Ile Val Ala Asp Cys Ile Leu Ala Gly Ala Tyr Val Leu Ser Gln Gly  
 1380 1385 1390  
 Arg Cys Lys Lys Cys Trp Gly Ser Cys Ile Arg Thr Ala Pro Asn Glu  
 1395 1400 1405  
 Val Ala Phe Asn Val Phe Pro Phe Thr Arg Ala Thr Arg Ser Ser Leu  
 1410 1415 1420  
 Ile Asp Leu Cys Asp Arg Phe Cys Ala Pro Lys Gly Met Asp Pro Ile  
 1425 1430 1435 1440  
 Phe Leu Ala Thr Gly Trp Arg Gly Cys Trp Ala Gly Arg Ser Pro Ile  
 1445 1450 1455  
 Glu Gln Pro Ser Glu Lys Pro Ile Ala Phe Ala Gln Leu Asp Glu Lys  
 1460 1465 1470  
 Lys Ile Thr Ala Arg Thr Val Val Ala Gln Pro Tyr Asp Pro Asn Gln  
 1475 1480 1485  
 Ala Val Lys Cys Leu Arg Val Leu Gln Ser Gly Gly Arg Trp Trp Leu  
 1490 1495 1500  
 Ser Gly Pro Lys Ser Gly Gln Gly Phe Arg Cys Ser Ile Pro Ser Pro  
 1505 1510 1515 1520  
 Phe Phe Pro Thr Gly Val Lys Val Asp Pro Asp Cys Arg Val Val Val  
 1525 1530 1535  
 Asp Pro Asp Thr Phe Thr Ala Ala Leu Arg Ser Gly Tyr Ser Thr Thr  
 1540 1545 1550  
 Asn Leu Val Leu Gly Val Gly Asp Phe Ala Gln Leu Asn Gly Leu Lys  
 1555 1560 1565  
 Ile Arg Gln Ile Ser Lys Pro Ser Gly Gly Gly Pro His Leu Met Ala  
 1570 1575 1580  
 Ala Leu His Val Ala Cys Ser Met Ala Leu His Met Leu Ala Gly Ile  
 1585 1590 1595 1600  
 Tyr Val Thr Ala Val Gly Ser Cys Gly Thr Gly Thr Asn Asp Pro Trp  
 1605 1610 1615  
 Cys Ala Asn Pro Phe Ala Val Pro Gly Tyr Gly Pro Gly Ser Leu Cys  
 1620 1625 1630  
 Thr Ser Arg Leu Cys Ile Ser Gln His Gly Leu Thr Leu Pro Leu Thr  
 1635 1640 1645

Ala Leu Val Ala Gly Phe Gly Ile Gln Glu Ile Ala Leu Val Val Leu  
 1650 1655 1660

Ile Phe Val Ser Ile Gly Gly Met Ala His Arg Leu Ser Cys Lys Ala  
 1665 1670 1675 1680

Asp Met Leu Cys Val Leu Leu Ala Ile Ala Ser Tyr Val Trp Val Pro  
 1685 1690 1695

Leu Thr Trp Leu Leu Cys Val Phe Pro Cys Trp Leu Arg Cys Phe Ser  
 1700 1705 1710

Leu His Pro Leu Thr Ile Leu Trp Leu Val Phe Phe Leu Ile Ser Val  
 1715 1720 1725

Asn Met Pro Ser Gly Ile Leu Ala Met Val Leu Leu Val Ser Leu Trp  
 1730 1735 1740

Leu Leu Gly Arg Tyr Thr Asn Val Ala Gly Leu Val Thr Pro Tyr Asp  
 1745 1750 1755 1760

Ile His His Tyr Thr Ser Gly Pro Arg Gly Val Ala Ala Leu Ala Thr  
 1765 1770 1775

Ala Pro Asp Gly Thr Tyr Leu Ala Ala Val Arg Arg Ala Ala Leu Thr  
 1780 1785 1790

Gly Arg Thr Met Leu Phe Thr Pro Ser Gln Leu Gly Ser Leu Leu Glu  
 1795 1800 1805

Gly Ala Phe Arg Thr Arg Lys Pro Ser Leu Asn Thr Val Asn Val Ile  
 1810 1815 1820

Gly Ser Ser Met Gly Ser Gly Gly Val Phe Thr Ile Asp Gly Lys Val  
 1825 1830 1835 1840

Lys Cys Val Thr Ala Ala His Val Leu Thr Gly Asn Ser Ala Arg Val  
 1845 1850 1855

Ser Gly Val Gly Phe Asn Gln Met Leu Asp Phe Asp Val Lys Gly Asp  
 1860 1865 1870

Phe Ala Ile Ala Asp Cys Pro Asn Trp Gln Gly Ala Ala Pro Lys Thr  
 1875 1880 1885

Gln Phe Cys Thr Asp Gly Trp Thr Gly Arg Ala Tyr Trp Leu Thr Ser  
 1890 1895 1900

Ser Gly Val Glu Pro Gly Val Ile Gly Lys Gly Phe Ala Phe Cys Phe  
 1905 1910 1915 1920

Thr Ala Cys Gly Asp Ser Gly Ser Pro Val Ile Thr Glu Ala Gly Glu  
 1925 1930 1935

Leu Val Gly Val His Thr Gly Ser Asn Lys Gln Gly Gly Gly Ile Val  
 1940 1945 1950

Thr	Arg	Pro	Ser	Gly	Gln	Phe	Cys	Asn	Val	Ala	Pro	Ile	Lys	Leu	Ser	1955	1960	1965	
Glu	Leu	Ser	Glu	Phe	Phe	Ala	Gly	Pro	Lys	Val	Pro	Leu	Gly	Asp	Val	1970	1975	1980	
Lys	Val	Gly	Ser	His	Ile	Ile	Lys	Asp	Ile	Ser	Glu	Val	Pro	Ser	Asp	1985	1990	1995	2000
Leu	Cys	Ala	Leu	Leu	Ala	Ala	Lys	Pro	Glu	Leu	Glu	Gly	Gly	Leu	Ser	2005	2010	2015	
Thr	Val	Gln	Leu	Leu	Cys	Val	Phe	Phe	Leu	Leu	Trp	Arg	Met	Met	Gly	2020	2025	2030	
His	Ala	Trp	Thr	Pro	Leu	Val	Ala	Val	Ser	Phe	Phe	Ile	Leu	Asn	Glu	2035	2040	2045	
Val	Leu	Pro	Ala	Val	Leu	Val	Arg	Ser	Val	Phe	Ser	Phe	Gly	Met	Phe	2050	2055	2060	
Val	Leu	Ser	Trp	Leu	Thr	Pro	Trp	Ser	Ala	Gln	Val	Leu	Met	Ile	Arg	2065	2070	2075	2080
Leu	Leu	Thr	Ala	Ala	Leu	Asn	Arg	Asn	Arg	Trp	Ser	Leu	Ala	Phe	Phe	2085	2090	2095	
Ser	Leu	Gly	Ala	Val	Thr	Gly	Phe	Val	Ala	Asp	Leu	Ala	Ala	Thr	Gln	2100	2105	2110	
Gly	His	Pro	Leu	Gln	Ala	Val	Met	Asn	Leu	Ser	Thr	Tyr	Ala	Phe	Leu	2115	2120	2125	
Pro	Arg	Met	Met	Val	Val	Thr	Ser	Pro	Val	Pro	Val	Ile	Thr	Cys	Gly	2130	2135	2140	
Val	Val	His	Leu	Leu	Ala	Ile	Ile	Leu	Tyr	Leu	Phe	Lys	Tyr	Arg	Gly	2145	2150	2155	2160
Pro	His	His	Ile	Leu	Val	Gly	Asp	Gly	Val	Phe	Ser	Ala	Ala	Phe	Phe	2165	2170	2175	
Leu	Arg	Tyr	Phe	Ala	Glu	Gly	Lys	Leu	Arg	Glu	Gly	Val	Ser	Gln	Ser	2180	2185	2190	
Cys	Gly	Met	Asn	His	Glu	Ser	Leu	Thr	Gly	Ala	Leu	Ala	Met	Arg	Leu	2195	2200	2205	
Asn	Asp	Glu	Asp	Leu	Asp	Phe	Leu	Met	Lys	Trp	Thr	Asp	Phe	Lys	Cys	2210	2215	2220	
Phe	Val	Ser	Ala	Ser	Asn	Met	Arg	Asn	Ala	Ala	Gly	Gln	Phe	Ile	Glu	2225	2230	2235	2240
Ala	Ala	Tyr	Ala	Lys	Ala	Leu	Arg	Val	Glu	Leu	Ala	Gln	Leu	Val	Gln	2245	2250	2255	



Val Asp Lys Val Arg Gly Thr Leu Ala Lys Leu Glu Ala Phe Ala Asp  
 2260 2265 2270  
 Thr Val Ala Pro Gln Leu Ser Pro Gly Asp Ile Val Val Ala Leu Gly  
 2275 2280 2285  
 His Thr Pro Val Gly Ser Ile Phe Asp Leu Lys Val Gly Ser Thr Lys  
 2290 2295 2300  
 His Thr Leu Gln Ala Ile Glu Thr Arg Val Leu Ala Gly Ser Lys Met  
 2305 2310 2315 2320  
 Thr Val Ala Arg Val Val Asp Pro Thr Pro Thr Pro Pro Ala Pro  
 2325 2330 2335  
 Val Pro Ile Pro Leu Pro Pro Lys Val Leu Glu Asn Gly Pro Asn Ala  
 2340 2345 2350  
 Trp Gly Asp Glu Asp Arg Leu Asn Lys Lys Lys Arg Arg Arg Met Glu  
 2355 2360 2365  
 Ala Leu Gly Ile Tyr Val Met Gly Gly Lys Lys Tyr Gln Lys Phe Trp  
 2370 2375 2380  
 Asp Lys Asn Ser Gly Asp Val Phe Tyr Glu Glu Val His Asn Asn Thr  
 2385 2390 2395 2400  
 Asp Glu Trp Glu Cys Leu Arg Val Gly Asp Pro Ala Asp Phe Asp Pro  
 2405 2410 2415  
 Glu Lys Gly Thr Leu Cys Gly His Val Thr Ile Glu Asn Lys Ala Tyr  
 2420 2425 2430  
 His Val Tyr Thr Ser Pro Ser Gly Lys Lys Phe Leu Val Pro Val Asn  
 2435 2440 2445  
 Pro Glu Asn Gly Arg Val Gln Trp Glu Ala Ala Lys Leu Ser Val Glu  
 2450 2455 2460  
 Gln Ala Leu Gly Met Met Asn Val Asp Gly Glu Leu Thr Ala Lys Glu  
 2465 2470 2475 2480  
 Leu Glu Lys Leu Lys Arg Ile Ile Asp Lys Leu Gln Gly Leu Thr Lys  
 2485 2490 2495  
 Glu Gln Cys Leu Asn Cys  
 2500

<210> 2

<211> 1457

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 2

Leu Ala Ala Ser Asp Leu Thr Arg Cys Gly Arg Gly Gly Leu Val Val

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Thr Glu Thr Ala Val Lys Ile Val Lys Phe His Asn Arg Thr Phe Thr	20	25	30
Leu Gly Pro Val Asn Leu Lys Val Ala Ser Glu Val Glu Leu Lys Asp	35	40	45
Ala Val Glu His Asn Gln His Pro Val Ala Arg Pro Ile Asp Gly Gly	50	55	60
Val Val Leu Leu Arg Ser Ala Val Pro Ser Leu Ile Asp Val Leu Ile	65	70	75
Ser Gly Ala Asp Ala Ser Pro Lys Leu Leu Ala His His Gly Pro Gly	85	90	95
Asn Thr Gly Ile Asp Gly Thr Leu Trp Asp Phe Glu Ser Glu Ala Thr	100	105	110
Lys Glu Glu Val Ala Leu Ser Ala Gln Ile Ile Gln Ala Cys Asp Ile	115	120	125
Arg Arg Gly Asp Ala Pro Glu Ile Gly Leu Pro Tyr Lys Leu Tyr Pro	130	135	140
Val Arg Gly Asn Pro Glu Arg Val Lys Gly Val Leu Gln Asn Thr Arg	145	150	155
Phe Gly Asp Ile Pro Tyr Lys Thr Pro Ser Asp Thr Gly Ser Pro Val	165	170	175
His Ala Ala Ala Cys Leu Thr Pro Asn Ala Thr Pro Val Thr Asp Gly	180	185	190
Arg Ser Val Leu Ala Thr Thr Met Pro Pro Gly Phe Glu Leu Tyr Val	195	200	205
Pro Thr Ile Pro Ala Ser Val Leu Asp Tyr Leu Asp Ser Arg Pro Asp	210	215	220
Cys Pro Lys Gln Leu Thr Glu His Gly Cys Glu Asp Ala Ala Leu Lys	225	230	235
Asp Leu Ser Lys Tyr Asp Leu Ser Thr Gln Gly Phe Val Leu Pro Gly	245	250	255
Val Leu Arg Leu Val Arg Lys Tyr Leu Phe Ala His Val Gly Lys Cys	260	265	270
Pro Pro Val His Arg Pro Ser Thr Tyr Pro Ala Lys Asn Ser Met Ala	275	280	285
Gly Ile Asn Gly Asn Arg Phe Pro Thr Lys Asp Ile Gln Ser Val Pro	290	295	300
Glu Ile Asp Val Leu Cys Ala Gln Ala Val Arg Glu Asn Trp Gln Thr			

305					310					315					320
Val	Thr	Pro	Cys	Thr	Leu	Lys	Lys	Gln	Tyr	Cys	Gly	Lys	Lys	Lys	Thr
				325					330					335	
Arg	Thr	Ile	Leu	Gly	Thr	Asn	Asn	Phe	Ile	Ala	Leu	Ala	His	Arg	Ala
			340					345					350		
Val	Leu	Ser	Gly	Val	Thr	Gln	Gly	Phe	Met	Lys	Lys	Ala	Phe	Asn	Ser
		355					360					365			
Pro	Ile	Ala	Leu	Gly	Lys	Asn	Lys	Phe	Lys	Glu	Leu	Gln	Thr	Pro	Val
	370					375					380				
Leu	Gly	Arg	Cys	Leu	Glu	Ala	Asp	Leu	Ala	Ser	Cys	Asp	Arg	Ser	Thr
385					390					395					400
Pro	Ala	Ile	Val	Arg	Trp	Phe	Ala	Ala	Asn	Leu	Leu	Tyr	Glu	Leu	Ala
				405					410					415	
Cys	Ala	Glu	Glu	His	Leu	Pro	Ser	Tyr	Val	Leu	Asn	Cys	Cys	His	Asp
			420					425					430		
Leu	Leu	Val	Thr	Gln	Ser	Gly	Ala	Val	Thr	Lys	Arg	Gly	Gly	Leu	Ser
		435					440					445			
Ser	Gly	Asp	Pro	Ile	Thr	Ser	Val	Ser	Asn	Thr	Ile	Tyr	Ser	Leu	Val
	450					455					460				
Ile	Tyr	Ala	Gln	His	Met	Val	Leu	Ser	Tyr	Phe	Lys	Ser	Gly	His	Pro
465					470					475					480
His	Gly	Leu	Leu	Phe	Leu	Gln	Asp	Gln	Leu	Lys	Phe	Glu	Asp	Met	Leu
				485					490					495	
Lys	Val	Gln	Pro	Leu	Ile	Val	Tyr	Ser	Asp	Asp	Leu	Val	Leu	Tyr	Ala
			500					505					510		
Glu	Ser	Pro	Thr	Met	Pro	Asn	Tyr	His	Trp	Trp	Val	Glu	His	Leu	Asn
		515					520					525			
Leu	Met	Leu	Gly	Phe	Gln	Thr	Asp	Pro	Lys	Lys	Thr	Ala	Ile	Thr	Asp
	530					535					540				
Ser	Pro	Ser	Phe	Leu	Gly	Cys	Arg	Ile	Ile	Asn	Gly	Arg	Gln	Leu	Val
545					550					555					560
Pro	Asn	Arg	Asp	Arg	Ile	Leu	Ala	Ala	Leu	Ala	Tyr	His	Met	Lys	Ala
				565					570					575	
Ser	Asn	Val	Ser	Glu	Tyr	Tyr	Ala	Ser	Ala	Ala	Ala	Ile	Leu	Met	Asp
			580					585					590		
Ser	Cys	Ala	Cys	Leu	Glu	Tyr	Asp	Pro	Glu	Trp	Phe	Glu	Glu	Leu	Val
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Lys Met Met Leu Asp Phe Lys Glu Val Arg Leu Met Val Trp Lys Asp 1285 1290 1295		
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Asn

<210> 3

<211> 256

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 3

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 Val Ala His Phe Gln His Leu Ala Ala Ile Glu Ala Glu Thr Cys Lys  
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 Tyr Leu Ala Ser Arg Leu Pro Met Leu His Asn Leu Arg Met Thr Gly  
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 Ile Phe Pro Thr Pro Gly Ser Arg Pro Lys Leu His Asp Phe Gln Gln  
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<210> 4

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<213> Porcine reproductive and respiratory syndrome virus

<400> 4

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<210> 5

<211> 4374

<212> DNA

<213> Porcine reproductive and respiratory syndrome virus

<400> 5

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<210> 6

<211> 771

<212> DNA

<213> Porcine reproductive and respiratory syndrome virus

<400> 6

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